

10/573522

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SEQUENCE LISTING

<110> Nakamura, Yusuke  
Katagiri, Toyomasa  
Nakatsuru, Shuichi

<120> METHOD OF DIAGNOSING BREAST CANCER

<130> 082368-007400US

<150> PCT/JP2004/011741  
<151> 2004-08-10

<150> US 60/505,571  
<151> 2003-09-24

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1 5 . 10

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15 20 . 25 30

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35 40 . 45

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Phe Glu Pro Pro Gln Ile Arg Phe Leu Thr Pro Ile Tyr His Pro Asn  
65 70 . 75

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80 85 . 90

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Lys Gly Ala Trp Arg Pro Ser Leu Asn Ile Ala Thr Val Leu Thr Ser  
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gacataattt ttgtgtagtt tatttatctt gtacatatgt attttgaaat cttttaaacc		870	
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Pro Pro Gln Ile Arg Phe Leu Thr Pro Ile Tyr His Pro Asn Ile Asp			
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 Gly Ser Arg Arg Arg Gln Arg Pro Ser Val Gly Val Gln Ser Leu  
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 Arg Pro Gln Ser Pro Gln Leu Arg Gln Ser Asp Pro Gln Lys Arg Asn  
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 Met Leu Ala Lys Leu His Glu Glu Ile Glu His Leu Lys Arg Glu Asn  
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 aag gat ctc cat tac aag ctc ata atg aat cag aca tca cag aag aaa 346  
 Lys Asp Leu His Tyr Lys Leu Ile Met Asn Gln Thr Ser Gln Lys Lys  
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 gat ggc ccc tca gga aac cac ctt tcc agg gcc tct gct ccc ttg ggc 394  
 Asp Gly Pro Ser Gly Asn His Leu Ser Arg Ala Ser Ala Pro Leu Gly  
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180	185	190	
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Asp Ser Lys Ala Asp Val Ser Gln Lys Ala Asp Leu Glu Glu Glu Pro			
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Leu Leu His Asn Ser Lys Leu Asp Lys Val Pro Gly Val Gln Gly Gln			
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Ala Arg Lys Glu Lys Ala Glu Ala Ser Asn Ala Gly Ala Ala Cys Met			
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Gly Asn Ser Gln His Gln Gly Arg Gln Met Gly Ala Gly Ala His Pro			
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Pro Met Ile Leu Pro Leu Pro Leu Arg Lys Pro Thr Thr Leu Arg Gln			
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caa gag ctg cgg cac ctc aag tcc ctc ctg gaa ggg agc cag agg ccc			970
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295	300	305	
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His Phe Pro Lys Val Ser Thr Lys Ser Leu Ser Lys Lys Cys Leu Ser			
325	330	335	
cca cct gtg gcg gag cgt gcc atc ctg ccc gca ctg aag cag acc ccg			1114
Pro Pro Val Ala Glu Arg Ala Ile Leu Pro Ala Leu Lys Gln Thr Pro			
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Lys Asn Asn Phe Ala Glu Arg Gln Lys Arg Leu Gln Ala Met Gln Lys			
355	360	365	370
cg <sup>g</sup> cg <sup>c</sup> ct <sup>g</sup> cat cg <sup>c</sup> tca gt <sup>g</sup> ctt tga gccacccaa tctggtcagt			1209
Arg Arg Leu His Arg Ser Val Leu *			
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 65          70          75          80
Glu Asn Lys Asp Leu His Tyr Lys Leu Ile Met Asn Gln Thr Ser Gln
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Lys Lys Asp Gly Pro Ser Gly Asn His Leu Ser Arg Ala Ser Ala Pro
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115         120         125
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Arg Ser Pro Ala Asp Ser Leu Ser Met Ser Ser Phe Gln Ser Val Lys
165         170         175
Ser Ile Ser Asn Ser Gly Lys Ala Arg Pro Gln Pro Gly Ser Phe Asn
180         185         190
Lys Gln Asp Ser Lys Ala Asp Val Ser Gln Lys Ala Asp Leu Glu Glu
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210         215         220
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Cys Met Gly Asn Ser Gln His Gln Gly Arg Gln Met Gly Ala Gly Ala
245         250         255
His Pro Pro Met Ile Leu Pro Leu Pro Leu Arg Lys Pro Thr Thr Leu
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275         280         285
Gln Thr Gln Glu Leu Arg His Leu Lys Ser Leu Leu Glu Gly Ser Gln
290         295         300
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Leu Ser Pro Pro Val Ala Glu Arg Ala Ile Leu Pro Ala Leu Lys Gln
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 aagcgcgggt gaagcgcgc aacagctgcgc tgccggcccg gctgcggtca 240  
 gcaacgcgcc atg gac gca gag ctg gca gag gtc cgc gcc ttg caa gct 289  
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gag atc gcg gcc ctg cgg cga gcg tgc ttt gag gac cca ccc ggc ccc tgg 337  
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gaa gag aag tcc cga gtc caa aaa tct ttt caa gcc ata cac caa ttc 385  
 Glu Glu Lys Ser Arg Val Gln Lys Ser Phe Gln Ala Ile His Gln Phe  
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aat ttg gaa gga tgg aag tct tca aaa gat ctg aaa aat cag ctt gga 433  
 Asn Leu Glu Gly Trp Lys Ser Ser Lys Asp Leu Lys Asn Gln Leu Gly  
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cat tta gaa tca gaa ctt tca ttt cta agt acg ctt act ggc atc aat 481  
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ata aga aat cac tcc aag cag aca gaa gac cta aca agc act gag atg 529  
 Ile Arg Asn His Ser Lys Gln Thr Glu Asp Leu Thr Ser Thr Glu Met  
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aca gaa aag agt att aga aaa gtt cta cag aga cac aga tta tca gga 577  
 Thr Glu Lys Ser Ile Arg Lys Val Leu Gln Arg His Arg Leu Ser Gly  
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aat tgc cac atg gtt aca ttt caa ctt gaa ttt cag att ctg gaa att 625  
 Asn Cys His Met Val Thr Phe Gln Leu Glu Phe Gln Ile Leu Glu Ile  
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gaa gag aga aaa gat ctg ttc atg ttt ttc cga agc ctg cat ttt ttt 769  
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gcc ata gaa act gct cct ctc agc ttc cga acc ctg gta gga ctg ctt		1057	
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Thr Ala Pro Leu Ser Phe Arg Thr Leu Val Gly Leu Leu Gly Ile Glu			
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RT-PCR

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